	CRF Errors Corrected by the STIC Systems French
Nι	Imber: 8/486,3/3 CRF Processing Date: 72/29/9 Edited by:
	Changed a file from non-ASCII to ASCII Verified by: (STIC s
1	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of files page numbers throughout text; ☐ other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (errodue to a Patentin bug). Sequences corrected:
	Other:

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 16:39:40

INPUT SET: S8022.raw

This Raw Listing contains the General **Information Section and those Sequences** containing ERRORS.

```
mayed got
1
                                       SEQUENCE LISTING
 2
 3
    (1)
           General Information:
 4
 5
          (i) APPLICANT: Weiss, Samuel
 6
                         Reynolds, Brent A.
 7
                         Hammang, Joseph P.
 8
                         Baetge, Edward E.
 9
10
         (ii) TITLE OF INVENTION: In Vitro and In Vivo
    Proliferation and
11
                 Use of Multipotent Neural Stem Cells and their
12
13
    Progeny
14
15
        (iii) NUMBER OF SEQUENCES: 8
16
17
         (iv) CORRESPONDENCE ADDRESS:
18
               (A) ADDRESSEE: Flehr, Hohbach, Test, Albritton
19
     & Herbert
20
               (B) STREET: Four Embarcadero Center, Suite 3400
21
               (C) CITY: San Francisco
22
               (D) STATE: California
23
               (E) COUNTRY: United States
24
               (F) ZIP: 94111-4187
25
          (V) COMPUTER READABLE FORM:
26
27
               (A) MEDIUM TYPE: Floppy disk
28
               (B) COMPUTER: IBM PC compatible
29
               (C) OPERATING SYSTEM: PC-DOS/MS-DOS
30
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
31
32
         (vi) CURRENT APPLICATION DATA:
33
               (A) APPLICATION NUMBER: US 08/486,313
34
               (B) FILING DATE: 07-JUN-1995
35
               (C) CLASSIFICATION:
36
37
       (viii) ATTORNEY/AGENT INFORMATION:
               (A) NAME: Brunelle, Jan P.
38
39
               (B) REGISTRATION NUMBER: 35,081
40
               (C) REFERENCE/DOCKET NUMBER: A-61105-11/DJB/JPB
41
42
         (ix) TELECOMMUNICATION INFORMATION:
43
               (A) TELEPHONE: (415) 781-1989
44
               (B) TELEFAX: (415) 398-3249
45
```

(C) TELEX: 910 277299

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 16:39:43

INPUT SET: S8022.raw

46 47

ERRORED SEQUENCES FOLLOW:

```
48
            (2) INFORMATION FOR SEQ ID NO:1:
       49
       50
                 (i) SEQUENCE CHARACTERISTICS:
-->
       51
                       (A) LENGTH: 20 base pairs
       52
                       (B) TYPE: nucleic acid
       53
                       (C) STRANDEDNESS: unknown
       54
                       (D) TOPOLOGY: unknown
       55
       56
                (ii) MOLECULE TYPE: CDNA
       57
       58
       59
       60
                (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
       61
       62
            GAGATGCGAC CCTCAGGGAC
       63
       64
                          20 -
       65
       66
            (2) INFORMATION FOR SEQ ID NO:2:
       67
       68
                 (i) SEQUENCE CHARACTERISTICS:
       69
                       (A) LENGTH: 20 base pairs
       70
                       (B) TYPE: nucleic acid
       71
                      (C) STRANDEDNESS: unknown
       72
                       (D) TOPOLOGY: unknown
       73
                (ii) MOLECULE TYPE: cDNA
       74
       75
       76
       77
       78
       79
                (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
       80
            GTCCCTGAGG GTCGCATCTC
       81
       82
                         20 _
       83
       84
            (2) INFORMATION FOR SEQ ID NO:3:
       85
       86
                 (i) SEQUENCE CHARACTERISTICS:
       87
                      (A) LENGTH: 19 base pairs
       88
                      (B) TYPE: nucleic acid
       89
                      (C) STRANDEDNESS: unknown
       90
                      (D) TOPOLOGY: unknown
```

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 16:39:45

```
91
 92
          (ii) MOLECULE TYPE: cDNA
 93
 94
 95
 96
 97
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
 98
 99
     TAAATAAAAG ATGCCCTGG
100
                   19
101
102
      (2) INFORMATION FOR SEQ ID NO:4:
103
104
           (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 19 base pairs
105
106
                (B) TYPE: nucleic acid
107
                (C) STRANDEDNESS: unknown
108
                (D) TOPOLOGY: unknown
109
110
          (ii) MOLECULE TYPE: cDNA
111
112
113
114
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
115
116
117
     CCAGGGCATC TTTTATTTA
118
119
      (2) INFORMATION FOR SEQ ID NO:5:
120
121
122
           (i) SEQUENCE CHARACTERISTICS:
123
                (A) LENGTH: 20 base pairs
124
                (B) TYPE: nucleic acid
125
                (C) STRANDEDNESS: unknown
                (D) TOPOLOGY: unknown
126
127
          (ii) MOLECULE TYPE: cDNA
128
129
130
131
132
133
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
134
135
      GAACTGGGAT GTGGGGCTGG
136
                   20
137
138
      (2) INFORMATION FOR SEQ ID NO:6:
139
140
           (i) SEQUENCE CHARACTERISTICS:
```

RAW SEQUENCE LISTING PAGE: 4 DATE: 12/29/95 PATENT APPLICATION US/08/486,313 TIME: 16:39:48 INPUT SET: S8022, raw (A) LENGTH: 20 base pairs --> (B) TYPE: nucleic acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: unknown (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6: CCAGCCCCAC ATCCCAGTTC (2) INFORMATION FOR SEQ ID NO:7: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: unknown (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7: GCCAGCGGCA TCACCTCG (2) INFORMATION FOR SEQ ID NO:8: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: unknown (ii) MOLECULE TYPE: cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

CGAGGTGATG CCGCTGGC

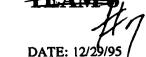
SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/486,313*

DATE: 12/29/95 TIME: 16:39:50

Line	Error	Original Text
51	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
69	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
87	Entered (19) and Calc. Seq. Length (0) differ	(A) LENGTH: 19 base pairs
105	Entered (19) and Calc. Seq. Length (0) differ	(A) LENGTH: 19 base pairs
123	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
141	Entered (20) and Calc. Seq. Length (0) differ	(A) LENGTH: 20 base pairs
159	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs
177	Entered (18) and Calc. Seq. Length (0) differ	(A) LENGTH: 18 base pairs

46

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313



DATE: 12/29/95 TIME: 17:07:25

INPUT SET: S8022.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

```
SEQUENCE LISTING
 1
 2
 3
    (1)
            General Information:
 5
          (i) APPLICANT: Weiss, Samuel
 6
                         Reynolds, Brent A.
                         Hammang, Joseph P.
 7
 8
                         Baetge, Edward E.
 9
         (ii) TITLE OF INVENTION: In Vitro and In Vivo Proliferation
10
              and Use of Multipotent Neural Stem Cells and their
11
12
              Progeny
13
        (iii) NUMBER OF SEQUENCES: 8
14
15
         (iv) CORRESPONDENCE ADDRESS:
16
               (A) ADDRESSEE: Flehr, Hohbach, Test, Albritton
17
                               & Herbert
18
               (B) STREET: Four Embarcadero Center, Suite 3400
19
20
               (C) CITY: San Francisco
               (D) STATE: California
21
               (E) COUNTRY: United States
22
23
               (F) ZIP: 94111-4187
24
25
          (v) COMPUTER READABLE FORM:
               (A) MEDIUM TYPE: Floppy disk
26
27
               (B) COMPUTER: IBM PC compatible
28
               (C) OPERATING SYSTEM: PC-DOS/MS-DOS
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
29
30
31
         (vi) CURRENT APPLICATION DATA:
               (A) APPLICATION NUMBER: US 08/486,313
32
33
               (B) FILING DATE: 07-JUN-1995
               (C) CLASSIFICATION:
34
35
       (viii) ATTORNEY/AGENT INFORMATION:
36
37
               (A) NAME: Brunelle, Jan P.
               (B) REGISTRATION NUMBER: 35,081
38
39
               (C) REFERENCE/DOCKET NUMBER: A-61105-11/DJB/JPB
40
         (ix) TELECOMMUNICATION INFORMATION:
41
               (A) TELEPHONE: (415) 781-1989
42
43
               (B) TELEFAX: (415) 398-3249
               (C) TELEX: 910 277299
44
45
```



RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 17:07:28

			1141 O1 3E1. 30022.74W
47 48	(2)	INFORMATION FOR SEQ ID NO:1:	
49		(i) SEQUENCE CHARACTERISTICS:	,
50		(A) LENGTH: 20 base pairs	
-			
51		(B) TYPE: nucleic acid	
52		(C) STRANDEDNESS: unknown	
53		(D) TOPOLOGY: unknown	•
54			
55		(ii) MOLECULE TYPE: cDNA	
56			
57			
58			
59			
60		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
61			
62	GAG	ATGCGAC CCTCAGGGAC	2
63			
64	(2)	INFORMATION FOR SEQ ID NO:2:	
65	` '		
66		(i) SEQUENCE CHARACTERISTICS:	
67		(A) LENGTH: 20 base pairs	
68		(B) TYPE: nucleic acid	
69		(C) STRANDEDNESS: unknown	
70		(D) TOPOLOGY: unknown	
71			
72		(ii) MOLECULE TYPE: cDNA	
73			
74			
75			
76			•
77		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
78		(iii) bagoanoa babonii iion. bag ib no.2.	
79	GTC(CCTGAGG GTCGCATCTC	20
80	G1C.	cerdade diederiere	20
81	(2)	TNEODNAMION FOR CHO ID NO. 2.	
82	(2)	INFORMATION FOR SEQ ID NO:3:	
83		(i) SEQUENCE CHARACTERISTICS:	·
84		(A) LENGTH: 19 base pairs	
85		(B) TYPE: nucleic acid	
86		(C) STRANDEDNESS: unknown	•
87		(D) TOPOLOGY: unknown	
88			
89		(ii) MOLECULE TYPE: cDNA	
90			
91			
92			
93			
94		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
95			
96	TAA	ATAAAAG ATGCCCTGG	19
97			
98	(2)	INFORMATION FOR SEQ ID NO:4:	
99	•		

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 17:07:31

		•	11/12 O = D=1/ D004=;· W.
100 101	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs	
102		(B) TYPE: nucleic acid	
103		(C) STRANDEDNESS: unknown	
		(D) TOPOLOGY: unknown	
104		(D) TOPOLOGY: Unknown	
105			
106	(ii) MOLECULE TYPE: cDNA	•
107			
108			
109			
110			
111	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	•
112	•	,	
113	CCAGGGC	ATC TTTTATTTA	~ 19
114	,		
115	/2\ TNE	ORMATION FOR SEQ ID NO:5:	
	(Z) INE	ORMATION FOR SEQ ID NO:5:	
116			
117	(1) SEQUENCE CHARACTERISTICS:	
118		(A) LENGTH: 20 base pairs	
119		(B) TYPE: nucleic acid	
120		(C) STRANDEDNESS: unknown	
121		(D) TOPOLOGY: unknown	
122			
123	(ii) MOLECULE TYPE: cDNA	
124			
125			
126			
127			
128	/xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
129	,	, and an entire result which is a second of the second of	
130	GAACTGG	GAT GTGGGGCTGG	20
131	GAAC 100	dar didddcidd	20
132	/2\ TNE	DODUMETON FOR GEO ID NO.C.	
	(Z) INE	ORMATION FOR SEQ ID NO:6:	
133			
134	(1) SEQUENCE CHARACTERISTICS:	
135		(A) LENGTH: 20 base pairs	
136		(B) TYPE: nucleic acid	
137		(C) STRANDEDNESS: unknown	
138		(D) TOPOLOGY: unknown	
139			
140	(ii	.) MOLECULE TYPE: cDNA	
141			•
142			
143		•	
144			
145	/yri) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
146	(24.1	.,E amountilion, and in no	
147	CCAGCCC	CCAC ATCCCAGTTC	20
	CCAGCCC	CAC AICCOAGIIC	20
148	(2)	PODMARION HOD CHO ID NO I	
149	(Z) INE	FORMATION FOR SEQ ID NO:7:	
150			
151	(i) SEQUENCE CHARACTERISTICS:	•
152		(A) LENGTH: 18 base pairs	

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 17:07:33

		INPUI SEI: Souzz.raw
153	(B) TYPE: nucleic acid	
154	(C) STRANDEDNESS: unknown	
155	(D) TOPOLOGY: unknown	
156		
157	(ii) MOLECULE TYPE: cDNA	
158	•	
159		
160	•	
161		
162	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
163		
164	GCCAGCGGCA TCACCTCG	18
165		
166	(2) INFORMATION FOR SEQ ID NO:8:	
167	(-,	
168	(i) SEQUENCE CHARACTERISTICS:	
169	(A) LENGTH: 18 base pairs	
170	(B) TYPE: nucleic acid	
171	(C) STRANDEDNESS: unknown	
172	(D) TOPOLOGY: unknown	
173	,	
174	(ii) MOLECULE TYPE: cDNA	
175	• •	
176		
177		
178		
179	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
180	· · · · ·	
181	CGAGGTGATG CCGCTGGC	18
182		
183		

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 17:07:35

INPUT SET: S8022.raw

***** PREVIOUSLY ERRORED SEQUENCES - EDITED *****

47	(2) INFORMATION FOR SEQ ID NO:1:	•
48		
49	(i) SEQUENCE CHARACTERISTICS:	
50	(A) LENGTH: 20 base pairs	
51	(B) TYPE: nucleic acid	
52	(C) STRANDEDNESS: unknown	
53	(D) TOPOLOGY: unknown	
54		
55	(ii) MOLECULE TYPE: cDNA	
56		
57		
58		
59		
60	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	•
61		
62	GAGATGCGAC CCTCAGGGAC	20
63		
64	(2) INFORMATION FOR SEQ ID NO:2:	
65 66	/ i \ CHOURNON CULD CONTRACTOR	
	(i) SEQUENCE CHARACTERISTICS:	
67	(A) LENGTH: 20 base pairs	
68	(B) TYPE: nucleic acid	
69	(C) STRANDEDNESS: unknown	
70	(D) TOPOLOGY: unknown	
71		
72	(ii) MOLECULE TYPE: cDNA	
73		
74		
75		
76		
77	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
78		
79	GTCCCTGAGG GTCGCATCTC	20
80		
81	(2) INFORMATION FOR SEQ ID NO:3:	
82	/-/	
83	(i) SEQUENCE CHARACTERISTICS:	
84	(A) LENGTH: 19 base pairs	
85	(B) TYPE: nucleic acid	
86	(C) STRANDEDNESS: unknown	
87	·	
88	(D) TOPOLOGY: unknown	•
89	(ii) MOLEGULE MUDE, abas	•
90	(ii) MOLECULE TYPE: cDNA	
90 91		
91		

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 17:07:38

		INPUT SET: S8022.raw
92		
93	,	
94	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
95		*
96	TAAATAAAAG ATGCCCTGG	. 19
97		
98	(2) INFORMATION FOR SEQ ID NO:4:	
99		
100	(i) SEQUENCE CHARACTERISTICS:	
101	(A) LENGTH: 19 base pairs	
102	(B) TYPE: nucleic acid	
103	(C) STRANDEDNESS: unknown	
104	(D) TOPOLOGY: unknown	
105	(11) MATERITE MURE, CANA	
106	(ii) MOLECULE TYPE: cDNA	
107		
108 109		
110		
111	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
111	(XI) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
113	CCAGGGCATC TTTTATTTA	19
114	CONGGGGATO TITTATTIA	2,5
115	(2) INFORMATION FOR SEQ ID NO:5:	
116		
117	(i) SEQUENCE CHARACTERISTICS:	
118	(A) LENGTH: 20 base pairs	
119	(B) TYPE: nucleic acid	
120 121	(C) STRANDEDNESS: unknown (D) TOPOLOGY: unknown	
121	(D) TOPOLOGI: UNKNOWN	
123	(ii) MOLECULE TYPE: cDNA	
124	(II) MODECODE IIIE. CDAR	
125		
126		•
127		
128	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
129	, , _	
130	GAACTGGGAT GTGGGGCTGG	20
131		
132	(2) INFORMATION FOR SEQ ID NO:6:	
133	• • • • • • • • • • • • • • • • • • • •	
134	(i) SEQUENCE CHARACTERISTICS:	
135	(A) LENGTH: 20 base pairs	
136		
137	(C) STRANDEDNESS: unknown	
138	(D) TOPOLOGY: unknown	
139		
140	(ii) MOLECULE TYPE: cDNA	
141		

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 17:07:40

142 143		
143	•	
144		
145	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
146	CO1CCCC10 1 maga2 amma	
147	CCAGCCCCAC ATCCCAGTTC	20
148		
149	(2) INFORMATION FOR SEQ ID NO:7:	
150		
151	(i) SEQUENCE CHARACTERISTICS:	
152	(A) LENGTH: 18 base pairs	
153	(B) TYPE: nucleic acid	
154	(C) STRANDEDNESS: unknown	
155	(D) TOPOLOGY: unknown	
156	(11) 101 54115 5 511	
157	(ii) MOLECULE TYPE: cDNA	
158 159		
160		
161	,	
162	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
163	(XI) SECONNEL DESCRIPTION: SEC ID NO:/:	
164	GCCAGCGGCA TCACCTCG	18
165		
166	(2) INFORMATION FOR SEQ ID NO:8:	
167	(=, management cont == = = = = = = = = = = = = = = = = =	
168	(i) SEQUENCE CHARACTERISTICS:	
169	(A) LENGTH: 18 base pairs	
170	(B) TYPE: nucleic acid	
171	(C) STRANDEDNESS: unknown	
172	(D) TOPOLOGY: unknown	
173		
174	(ii) MOLECULE TYPE: cDNA	
175		
176		
177		
178	(-1) GEOVERNOR BEGGETTERS	
179	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	·
180 181	COLOGRADAMO, COCOMOGO	
181	CGAGGTGATG CCGCTGGC	18
182		

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/486,313

DATE: 12/29/95 TIME: 17:07:42

INPUT SET: S8022.raw

Line

Error

Original Text